

SCORE Search Results Details for Application 10622896 and Search Result us-10-622-896- 1.rge.

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This page gives you Search Results detail for the Application 10622896 and Search Result us-10-622-896-1.rge.

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GenCore version 5.1.9
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OM nucleic - nucleic search, using sw model

Run on: July 28, 2006, 16:31:03 ; Search time 10201 Seconds
(without alignments)
11515.676 Million cell updates/sec

Title: US-10-622-896-1

Perfect score: 1837

Sequence: 1 gactcactataggcggccg.....tctttgtttggtagtcttt 1837

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 6366136 seqs, 31973710525 residues

Total number of hits satisfying chosen parameters: 12732272

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : GenEmbl:*

1: gb_env:*

2: gb_pat:*

3: gb_ph:*

4: gb_pl:*

5: gb_pr:*

6: gb_ro:*

7: gb_sts:*

8: gb_sy:*

9: gb_un:*

10: gb_vi:*

11: gb_ov:*

12: gb_htg:*

13: gb_in:*

14: gb_om:*

15: gb_ba:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

%

Result No.	Score	Query Match	Length	DB ID	Description
1	1380.4	75.1	1726	5 AY147881	AY147881 Homo sapi
c 2	1368.8	74.5	2976	2 CQ412068	CQ412068 Sequence